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添付公開書類
国際調査報告書
補正書

(54) Title : GENE SIGNATURE

(54) 発明の名称 ジーン・シグナチャー

(57) Abstract

A 3'-directed cDNA library which accurately reflects the abundance ratio of mRNA in a cell has been prepared from various human tissues, and sequencing of the cDNAs contained in the library has been conducted to examine the incidence of each cDNA in each tissue. As each cDNA has expression information with each tissue corresponding to the mRNA concentration, these cDNAs are usable as a probe or primer for detecting cell anomaly or discriminating cells. The cloned gene can produce proteins utilizable as a medicine or the like.

AAGGGAACCT GCTTCTTTAC TCCAGAACTC TGTTCTTTAA AGACCAAGAT TACATTCTCA 180
ATTAGAAAAC TGCAATTTGG TTCCACCACA TCCTGACTAC TACCGTATAG TTTTCTCTAT 240
TCTTTCATTT CCCCCTTCCC CATTCCTTTA TTGTACATAA AGTAACTGGT ATATGTGCAC 300
AAA 303

配列番号：3165
配列の長さ：311
配列の型：核酸
トポロジー：直鎖状
クローン名：HUMGS03736
配列：
GATCTAGAGC CAGGCTGGTC AAGACAAGAA TTGGCTGGAA TAGGCTGCTC TTCCCCCATT 60
CCATCATGTG CTGTCCCCAC CCCTTTGGCC ACCTGGGCTG ACTGTGTCTT AATACCTCAA 120
GTGCAAGTAT ATAGGAGTAA GAAATNAACA ATGCCTGCCT CCTTATACTC ATGCCTACAT 180
TGTATGACAT CTAGTATGAA AGGGAAACAT TAAAGGAAAA CCCTTGTTTT GCTCTAAAAC 240
CTGAGGACGG TAAACACTGA GAGTAACCTG GTGCTTGGTT TGAAGTAAAA CACAAATACT 300
TCCCTTTTAA A 311

配列番号：3166
配列の長さ：309
配列の型：核酸
トポロジー：直鎖状
クローン名：HUMGS03737
配列：
GATCAAGGGC AATGCCAATG AACATCGGCA TGGATTATAA TTATGCCCTC CTGGAACTCA 60
AAAAGCCCCA CAAGAGAAAA TTTATGAAGA TTGGGGTGAG CCCTCCTGCT AAGCAGCTGC 120
CAGGGGGCAG AATTCACTTC TCTGGTTATN ACAATNNCCN NCCAGGCAAT TTGGTGTATC 180
GCTTCTGTGA CGTCAAAGAC GAGACCTATG ACTTGCTCTA CCAGCAATGC GATGCCCAGC 240
CAGGGGCCAG CGGGTCTGGG GTCTATGTGA GGATGTGGAA GAGACAGCAG CAGAAGTGGG 300
AGCGAAAAN 309

配列番号：3167
配列の長さ：309
配列の型：核酸
トポロジー：直鎖状
クローン名：HUMGS03738
配列：
GATCTTGATA TGTTTTAACA TTATCATGGC AGGGAAATAT ATAAAGAAGA AAAATATTTT 60
NACATTAAAC CTTTTCTAAA ANTTGTAAAT AGAAAAATAA TTTGGTTTTT NATCAAGANC 120
AACACTTATC GTTATGTATT GTGTTAGTTA TATTGCCAGT CTGTTGCGAC TGACTCAAAA 180
AGTTAAATGT TGCCACTGCT GANGATGATT ATGNGCATCG CANACTTTGT TTCTGNCCCA 240
TTTTGGCAGT TTTNATATAC TCCNTTAAGA TGTTGAATGT TACAGGTTAN TAAAGTTAAT 300
ACCTTTAAA 309

配列番号：3168

PSN_T22166 |GENESEQ-N| Human gene signature HUMGS03737

;ID T22166 standard; cDNA to mRNA; 309 BP.
;AC T22166;
;DT 08-AUG-1996 (first entry)
;DE Human gene signature HUMGS03737.
;KW Gene signature; messenger RNA; mRNA; relative abundance; frequency;
;KW human; cloning; mapping; non-biased library; diagnosis; detection;
;KW cell typing; abnormal cell function; ss.
;OS Homo sapiens.
;PN WO9514772-A1.
;PD 01-JUN-1995.
;PF 11-NOV-1994; J01916.
;PR 12-NOV-1993; JP-355504.
;PA (MATS/) MATSUBARA K.
;PA (OKUB/) OKUBO K.
;PI Matsubara K, Okubo K;
;DR WPI; 95-206931/27.
;PT Identifying gene signatures in 3'-directed human cDNA library - e.g.
;PT for diagnosis of abnormal cell function, by preparing cDNA that
;PT reflects relative abundance of corresp. mRNA in specific human
;PT tissues
;PS Claim 1; Page 1060; 2245pp; Japanese.
;CC A single-stranded DNA (or its complementary strand or the corresp.
;CC double-stranded DNA) which comprises one of the 7837 "GS" sequences
;CC given in T19001-T26837 and which is able to hybridise to part of
;CC human genomic DNA, cDNA or mRNA is claimed. The GS (Gene Signature)
;CC sequences were obtained from 3'-directed cDNA libraries prepared
;CC from various human tissues; synthesis of cDNA was initiated from the
;CC 3'-end of mRNA by using poly(T) as the sole primer. Since the 3'-
;CC untranslated sequence is unique to a particular mRNA species, almost
;CC all the 3'-oriented cDNAs hybridise with specific mRNAs. Each library
;CC is constructed so as to reflect accurately the relative abundance of
;CC different mRNAs in the particular tissue from which it was derived.
;CC The appearance frequency of a given GS in a cDNA library can be
;CC determined (esp. using primers and probes derived from the GS
;CC sequences) as a means of diagnosing abnormal cell function or for
;CC recognising different cell types.